OIPE

RAW SEQUENCE LISTING   DATE: 07/26/2001
PATENT APPLICATION: US/09/857,008A   TIME: 10:26:28

Input Set : **A:\PTO_VSK.txt**
Output Set: **N:\CRF3\07262001\I857008A.raw**

3 <110> APPLICANT: Herscovici, Jean
4   Hofland, Hans
5   Jacopin, Christophe
6   Scherman, Daniel
8 <120> TITLE OF INVENTION: NOVEL NUCLEIC ACID TRANSFERRING AGENTS, COMPOSITIONS
CONTAINING THEM AND
9   USES
11 <130> FILE REFERENCE: ST98046PC1-US
13 **<140> CURRENT APPLICATION NUMBER: US/09/857,008A**
14 **<141> CURRENT FILING DATE: 2001-07-13**
16 <150> PRIOR APPLICATION NUMBER: PCT/FR99/02995
17 <151> PRIOR FILING DATE: 1999-12-02
19 <150> PRIOR APPLICATION NUMBER: US 60/117,184
20 <151> PRIOR FILING DATE: 1999-01-26
22 <150> PRIOR APPLICATION NUMBER: FR 98/15,309
23 <151> PRIOR FILING DATE: 1998-12-03
25 <160> NUMBER OF SEQ ID NOS: 2
27 <170> SOFTWARE: PatentIn version 3.0
29 <210> SEQ ID NO: 1
30 <211> LENGTH: 9
31 <212> TYPE: PRT
32 <213> ORGANISM: none
34 <400> SEQUENCE: 1
36 Lys Thr Pro Lys Lys Ala Lys Lys Pro
37 1   5
39 <210> SEQ ID NO: 2
40 <211> LENGTH: 8
41 <212> TYPE: PRT
42 <213> ORGANISM: none
44 <400> SEQUENCE: 2
46 Ala Thr Pro Ala Lys Lys Ala Ala
47 1   5

VERIFICATION SUMMARY   DATE: 07/26/2001
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L:13 M:270 C: Current Application Number differs, Replaced Current Application Number
L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: 09/857,0084

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1 ____ Wrapped Nucleics
Wrapped Aminos

The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2 ____ Invalid Line Length

The rules require that a line not exceed 72 characters in length. This includes white spaces.

3 ____ Misaligned Amino
Numbering

The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use **space characters**, instead.

4 ____ Non-ASCII

The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**

5 ____ Variable Length

Sequence(s) ____ contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6 ____ PatentIn 2.0
"bug"

A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**

7 ____ Skipped Sequences
(OLD RULES)

Sequence(s) ____ missing. If intentional, please insert the following lines for **each** skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.

8 ____ Skipped Sequences
(NEW RULES)

Sequence(s) ____ missing. If **intentional**, please insert the following lines for **each** skipped sequence.
<210> sequence id number
<400> sequence id number
000

9 ____ Use of n's or Xaa's
(NEW RULES)

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10 ____ Invalid <213>
Response

Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence

11 ____ Use of <220>

Sequence(s) ____ missing the <220> "Feature" and associated numeric identifiers and responses Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No 104, pp 29631-32)   (Sec. 1.823 of Sequence Rules)

12 ____ PatentIn 2.0
"bug"

Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13 ____ Misuse of n

n can only be used to represent a single **nucleotide** in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.